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1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/721,904B

DATE: 03/20/2003 P.6

TIME: 16:03:44

Input Set : A:\seqlist.asc.txt

Output Set: N:\CRF4\03202003\I721904B.raw

3 <110> APPLICANT: JULIUS, Michael H.
4 FILIPP, Dominik
6 <120> TITLE OF INVENTION: THE INDUCTION OF ANTIBIOTIC PROTEINS AND PEPTIDES BY
7 LAIT/sCD14-PROTEIN
9 <130> FILE REFERENCE: 47841/00063
11 <140> CURRENT APPLICATION NUMBER: US 09/721,904B
12 <141> CURRENT FILING DATE: 2000-11-27
14 <150> PRIOR APPLICATION NUMBER: PCT/CA99/00482
15 <151> PRIOR FILING DATE: 1999-05-27
17 <150> PRIOR APPLICATION NUMBER: US 60/086,884
18 <151> PRIOR FILING DATE: 1998-05-27
20 <160> NUMBER OF SEQ ID NOS: 11
22 <170> SOFTWARE: Wordperfect 9.0
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25 <211> LENGTH: 1122
26 <212> TYPE: DNA
27 <213> ORGANISM: bovine
29 <400> SEQUENCE: 1
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32 gacacaacag aaccctgcga gctggacgac gacgatttc gttgtgtctg caacttcacg 120
34 gatccgaaggc ctgactggc tagcgccgtt cagtgtatgg ttgcgcgtcga ggtggagatc 180
36 agtgccggcg gccgcggcct ggaacagttt ctcaaggag ccgcacaccaa cccgaagcag 240
38 tatgctgaca caatcaaggc tctgcgcgtt cggcgaactca agctgggcgc tgcacaggtt 300
40 cctgctcagc ttctggcgcg cgttctgcgc gcgctgggt actctcgctt caaggaactg 360
42 acgcttgagg acctggaggt aaccggccca acgccccca cgcctctgga agccgctggg 420
44 cctgcgtca ccacccttag tctgcgttaac gtatcggtt caacaggagg tgcctggctc 480
46 ggcgaactgc agcagtggct caagcctggg ctcagggtgc tgaacattgc ccaagcacac 540
48 tcgcttgccc ttccgtgcgc agggctctcc accttcgagg cgctcaccac cctagacactg 600
50 tctgacaatc ccagtctcgg cgacacgggg ctgatggcag ctctctgtcc gaacaagttc 660
52 cccggccctcc aatatctagc gctacgcaac gcggggatgg agacgcccgg cggcggtgtc 720
54 gccgcgtgg cggcagcgg ggtgcagccc caaaggctgg acctcagcca caactcgctg 780
56 cccgtcaccgc ccccggtgc tacccgatgt gtctggccca gtgcactaaag gtctctcaat 840
58 ttgtcggtcg ctgggttgg gcaagtgcct aaggactgc cccctaagct cagcgtgtt 900
60 gatctcagct gcaacaagct aaggcaggag cccgcggcgg acgagctgcc cgaggtaat 960
62 gacctgactc tggacggaaa tccctttctg gaccctggag ccctccagca caaaaatgac 1020
64 cccatgtatct cccgcgtggt cccagcctgt ggcgttctg cttgaccat ggggggtgtca 1080
66 ggagccctgg cgctgcttca aggagccga ggcttcgcgt aa 1122
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70 <211> LENGTH: 1128
71 <212> TYPE: DNA
72 <213> ORGANISM: human
74 <400> SEQUENCE: 2
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77	acgccagaac	cttgtgagct	ggacgatgaa	gattccgct	gcgtctgcaa	cttcccgaa	120	
79	cctcagcccg	actggtccga	agccttccag	tgtgtctcg	cagttaggt	ggagatccat	180	
81	gccggcggtc	tcaacctaga	gccgtttcta	aagcgcgtcg	atgcggacgc	cgaccgcgg	240	
83	cagtatgctg	acacggtaa	ggctctccgc	gtgcggcggc	tcacagtggg	agccgcacag	300	
85	gttcctgctc	agctactggt	aggcgcctg	cgtgtctag	cgtactccc	cctcaaggaa	360	
87	ctgacgctcg	aggaccta	aataaccggc	accatgcctc	cgctgcctc	ggaagccaca	420	
89	ggacttgcac	tttccagctt	gcmcctacgc	aacgtgtcg	ggcgacagg	gcgttcttg	480	
91	ctcgccgagc	tgcagcagt	gctcaagcca	ggcctaagg	tactgagcat	tgcccaagca	540	
93	cactcgctg	cctttcctg	cgaacaggtt	cgcgccttcc	cggcccttac	cagctagac	600	
95	ctgtctgaca	atcctggact	gggcgaacgc	ggactgtatgg	cggctctctg	tccccacaag	660	
97	ttcccgccca	tccagaatct	agegctgcgc	aacacaggaa	tggagacggc	cacaggcgtg	720	
99	tgcgcgcac	ttggcggcggc	aggtgtcg	ccccacagcc	tagacctca	ccacaactcg	780	
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103	ctcaatctgt	cgttcgctgg	gctggAACAG	gtgcctaaag	gactgccagc	caagctcaga	900	
105	gtgctcgatc	tcagctgca	cagactgaac	aggcgcggc	agcctgaoga	gctgcccgg	960	
107	gtggataacc	tgacactgga	cggaaatccc	ttcctggtcc	ctggaaactgc	cctcccccac	1020	
109	gagggctcaa	tgaactccgg	cgtggccca	gcctgtgcac	gttcgaccct	gtcggtgggg	1080	
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115	<211>	LENGTH:	1101					
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117	<213>	ORGANISM:	murine					
119	<400>	SEQUENCE:	3					
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122	gagccctgcg	agctagacga	ggaaagtgt	tcctgcaact	tctcagatcc	gaagccagat		120
124	tggccagcg	ctttcaattt	tttggggggc	gcagatgtgg	aattgtacgg	cgccggccgc		180
126	agcctggaa	atcctctaa	gcgtgtggac	acggaaggc	atctggggca	gttcaactgat		240
128	attatcaagt	ctctgtcctt	aaagcggctt	acgggtcg	ccgcgcggat	tccttagtcgg		300
130	attctattcg	gagccctgcg	tgtgtcg	atttccggcc	tccaggaact	gactcttggaa		360
132	aatctcgagg	taaccggcac	cgcgcggcca	ccgcctctgg	aagccacccgg	acccgatctc		420
134	aacatcttga	acctccgca	cgtgtcg	gcaacaagg	atgcctggct	cgcagaactg		480
136	cagcgtggc	taaagcctgg	actcaaggta	ctgagtattt	cccaagcaca	ctcaactcaac		540
138	ttttcctgcg	aacaggtccg	cgtttccct	gccctctcca	ccttagac	gtctgacaat		600
140	cctgaattgg	gcgagagagg	actgatctca	gccctctgc	ccctcaagtt	ccgcaccctc		660
142	caagtttag	cgctcgtaa	cgcggggat	gagacgccc	gcggcgtgt	ctctgcgt		720
144	gccgcagcaa	gggtacagct	gcaaggacta	gacccttagt	acaatttact	gcgggatgt		780
146	gcaggcgtc	cgagttgtga	ctggccca	cagctaaact	cgctcaatct	gtctttact		840
148	gggctgaagc	aggtaactaa	aggcgtccca	gccaagctca	gcgtgtcg	tctcagttac		900
150	aacaggctgg	ataggaaccc	tagcccagat	gagtcggccc	aagtggggaa	cctgtcaact		960
152	aaaggaaatc	ccttttgg	ctctgaatcc	cactcgaga	agtttaactc	tggcgtagt		1020
154	accgcggag	ctccatcatc	ccaagcagt	gcctgtcg	gaactctggc	tttgctctta		1080
156	ggagatcgcc	tctttgttta	a					1101
159	<210>	SEQ ID NO:	4					
160	<211>	LENGTH:	373					
161	<212>	TYPE:	PRT					
162	<213>	ORGANISM:	bovine					
164	<400>	SEQUENCE:	4					
165	Met Val Cys Val Pro Tyr Leu Leu Leu Leu Leu Pro Ser Leu Leu							
166	1	5		10		15		

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```

168 Arg Val Ser Ala Asp Thr Thr Glu Pro Cys Glu Leu Asp Asp Asp Asp
169      20          25          30
171 Phe Arg Cys Val Cys Asn Phe Thr Asp Pro Lys Pro Asp Trp Ser Ser
172      35          40          45
174 Ala Val Gln Cys Met Val Ala Val Glu Val Glu Ile Ser Ala Gly Gly
175      50          55          60
177 Arg Ser Leu Glu Gln Phe Leu Lys Gly Ala Asp Thr Asn Pro Lys Gln
178 65      70          75          80
180 Tyr Ala Asp Thr Ile Lys Ala Leu Arg Val Arg Arg Leu Lys Leu Gly
181      85          90          95
183 Ala Ala Gln Val Pro Ala Gln Leu Leu Val Ala Val Leu Arg Ala Leu
184      100         105         110
186 Gly Tyr Ser Arg Leu Lys Glu Leu Thr Leu Glu Asp Leu Glu Val Thr
187      115         120         125
189 Gly Pro Thr Pro Pro Thr Pro Leu Glu Ala Ala Gly Pro Ala Leu Thr
190      130         135         140
192 Thr Leu Ser Leu Arg Asn Val Ser Trp Thr Thr Gly Gly Ala Trp Leu
193 145      150         155         160
195 Gly Glu Leu Gln Gln Trp Leu Lys Pro Gly Leu Arg Val Leu Asn Ile
196      165         170         175
198 Ala Gln Ala His Ser Leu Ala Phe Pro Cys Ala Gly Leu Ser Thr Phe
199      180         185         190
201 Glu Ala Leu Thr Thr Leu Asp Leu Ser Asp Asn Pro Ser Leu Gly Asp
202      195         200         205
204 Thr Gly Leu Met Ala Ala Leu Cys Pro Asn Lys Phe Pro Ala Leu Gln
205      210         215         220
207 Tyr Leu Ala Leu Arg Asn Ala Gly Met Glu Thr Pro Ser Gly Val Cys
208 225      230         235         240
210 Ala Ala Leu Ala Ala Ala Arg Val Gln Pro Gln Ser Leu Asp Leu Ser
211      245         250         255
213 His Asn Ser Leu Arg Val Thr Ala Pro Gly Ala Thr Arg Cys Val Trp
214      260         265         270
216 Pro Ser Ala Leu Arg Ser Leu Asn Leu Ser Phe Ala Gly Leu Glu Gln
217      275         280         285
219 Val Pro Lys Gly Leu Pro Pro Lys Leu Ser Val Leu Asp Leu Ser Cys
220      290         295         300
222 Asn Lys Leu Ser Arg Glu Pro Arg Arg Asp Glu Leu Pro Glu Val Asn
223 305      310         315         320
225 Asp Leu Thr Leu Asp Gly Asn Pro Phe Leu Asp Pro Gly Ala Leu Gln
226      325         330         335
228 His Gln Asn Asp Pro Met Ile Ser Gly Val Val Pro Ala Cys Ala Arg
229      340         345         350
231 Ser Ala Leu Thr Met Gly Val Ser Gly Ala Leu Ala Leu Leu Gln Gly
232      355         360         365
234 Ala Arg Gly Phe Ala
235      370
238 <210> SEQ ID NO: 5
239 <211> LENGTH: 375
240 <212> TYPE: PRT

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RAW SEQUENCE LISTING
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241 <213> ORGANISM: human
 243 <400> SEQUENCE: 5
 244 Met Glu Arg Ala Ser Cys Leu Leu Leu Leu Leu Pro Leu Val His
 245 1 5 10 15
 247 Val Ser Ala Thr Thr Pro Glu Pro Cys Glu Leu Asp Asp Glu Asp Phe
 248 20 25 30
 250 Arg Cys Val Cys Asn Phe Ser Glu Pro Gln Pro Asp Trp Ser Glu Ala
 251 35 40 45
 253 Phe Gln Cys Val Ser Ala Val Glu Val Glu Ile His Ala Gly Gly Leu
 254 50 55 60
 256 Asn Leu Glu Pro Phe Leu Lys Arg Val Asp Ala Asp Ala Asp Pro Arg
 257 65 70 75 80
 259 Gln Tyr Ala Asp Thr Val Lys Ala Leu Arg Val Arg Arg Leu Thr Val
 260 85 90 95
 262 Gly Ala Ala Gln Val Pro Ala Gln Leu Leu Val Gly Ala Leu Arg Val
 263 100 105 110
 265 Leu Ala Tyr Ser Arg Leu Lys Glu Leu Thr Leu Glu Asp Leu Lys Ile
 266 115 120 125
 268 Thr Gly Thr Met Pro Pro Leu Pro Leu Glu Ala Thr Gly Leu Ala Leu
 269 130 135 140
 271 Ser Ser Leu Arg Leu Arg Asn Val Ser Trp Ala Thr Gly Arg Ser Trp
 272 145 150 155 160
 274 Leu Ala Glu Leu Gln Gln Trp Leu Lys Pro Gly Leu Lys Val Leu Ser
 275 165 170 175
 277 Ile Ala Gln Ala His Ser Pro Ala Phe Ser Tyr Glu Gln Val Arg Ala
 278 180 185 190
 280 Phe Pro Ala Leu Thr Ser Leu Asp Leu Ser Asp Asn Pro Gly Leu Gly
 281 195 200 205
 283 Glu Arg Gly Leu Met Ala Ala Leu Cys Pro His Lys Phe Pro Ala Ile
 284 210 215 220
 286 Gln Asn Leu Ala Leu Arg Asn Thr Gly Met Glu Thr Pro Thr Gly Val
 287 225 230 235 240
 289 Cys Ala Ala Leu Ala Ala Gly Val Gln Pro His Ser Leu Asp Leu
 290 245 250 255
 292 Ser His Asn Ser Leu Arg Ala Thr Val Asn Pro Ser Ala Pro Arg Cys
 293 260 265 270
 295 Met Trp Ser Ser Ala Leu Asn Ser Leu Asn Leu Ser Phe Ala Gly Leu
 296 275 280 285
 298 Glu Gln Val Pro Lys Gly Leu Pro Ala Lys Leu Arg Val Leu Asp Leu
 299 290 295 300
 301 Ser Cys Asn Arg Leu Asn Arg Ala Pro Gln Pro Asp Glu Leu Pro Glu
 302 305 310 315 320
 304 Val Asp Asn Leu Thr Leu Asp Gly Asn Pro Phe Leu Val Pro Gly Thr
 305 325 330 335
 307 Ala Leu Pro His Glu Gly Ser Met Asn Ser Gly Val Val Pro Ala Cys
 308 340 345 350
 310 Ala Arg Ser Thr Leu Ser Val Gly Val Ser Gly Thr Leu Val Leu Leu
 311 355 360 365
 313 Gln Gly Ala Arg Gly Phe Ala

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Input Set : A:\seqlist.asc.txt
Output Set: N:\CRF4\03202003\I721904B.raw

314	370	375
317 <210>	SEQ ID NO: 6	
318 <211>	LENGTH: 366	
319 <212>	TYPE: PRT	
320 <213>	ORGANISM: murine	
322 <400>	SEQUENCE: 6	
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324 1	5	10 15
326	Pro Ala Pro Pro Glu Pro Cys Glu Leu Asp Glu Glu Ser Cys Ser Cys	
327	20	25 30
329	Asn Phe Ser Asp Pro Lys Pro Asp Trp Ser Ser Ala Phe Asn Cys Leu	
330	35	40 45
332	Gly Ala Ala Asp Val Glu Leu Tyr Gly Gly Arg Ser Leu Glu Tyr	
333	50	55 60
335	Leu Leu Lys Arg Val Asp Thr Glu Ala Asp Leu Gly Gln Phe Thr Asp	
336	65	70 75 80
338	Ile Ile Lys Ser Leu Ser Leu Lys Arg Leu Thr Val Arg Ala Ala Arg	
339	85	90 95
341	Ile Pro Ser Arg Ile Leu Phe Gly Ala Leu Arg Val Leu Gly Ile Ser	
342	100	105 110
344	Gly Leu Gln Glu Leu Thr Leu Glu Asn Leu Glu Val Thr Gly Thr Ala	
345	115	120 125
347	Pro Pro Pro Leu Leu Glu Ala Thr Gly Pro Asp Leu Asn Ile Leu Asn	
348	130	135 140
350	Leu Arg Asn Val Ser Trp Ala Thr Arg Asp Ala Trp Leu Ala Glu Leu	
351	145	150 155 160
353	Gln Gln Trp Leu Lys Pro Gly Leu Lys Val Leu Ser Ile Ala Gln Ala	
354	165	170 175
356	His Ser Leu Asn Phe Ser Cys Glu Gln Val Arg Val Phe Pro Ala Leu	
357	180	185 190
359	Ser Thr Leu Asp Leu Ser Asp Asn Pro Glu Leu Gly Glu Arg Gly Leu	
360	195	200 205
362	Ile Ser Ala Leu Cys Pro Leu Lys Phe Pro Thr Leu Gln Val Leu Ala	
363	210	215 220
365	Leu Arg Asn Ala Gly Met Glu Thr Pro Ser Gly Val Cys Ser Ala Leu	
366	225	230 235 240
368	Ala Ala Ala Arg Val Gln Leu Gln Gly Leu Asp Leu Ser His Asn Ser	
369	245	250 255
371	Leu Arg Asp Ala Ala Gly Ala Pro Ser Cys Asp Trp Pro Ser Gln Leu	
372	260	265 270
374	Asn Ser Leu Asn Leu Ser Phe Thr Gly Leu Lys Gln Val Pro Lys Gly	
375	275	280 285
377	Leu Pro Ala Lys Leu Ser Val Leu Asp Leu Ser Tyr Asn Arg Leu Asp	
378	290	295 300
380	Arg Asn Pro Ser Pro Asp Glu Leu Pro Gln Val Gly Asn Leu Ser Leu	
381	305	310 315 320
383	Lys Gly Asn Pro Phe Leu Asp Ser Glu Ser His Ser Glu Lys Phe Asn	
384	325	330 335
386	Ser Gly Val Val Thr Ala Gly Ala Pro Ser Ser Gln Ala Val Ala Leu	

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/20/2003
PATENT APPLICATION: US/09/721,904B TIME: 16:03:45

Input Set : A:\seqlist.asc.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; Xaa Pos. 14,265,266,267,269

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/721,904B

DATE: 03/20/2003
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L:120 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=3
L:414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
M:341 Repeated in SeqNo=7